

| | | |
|------|---|-----|
| NT1p | MGLHANSFLKDFLAGVAAAVSKTAVAPIERVKLLLVQHASKQISAEKQ | 50 |
| NT2p | MTLPAVSAFAKDFLAGGVAAALSKTAVAPIERVKLLLVQHASKQIADKQ | 50 |
| NT3p | MTEQATSAFAKDFLAGGIAAALSKTAVAPIERVKLLLVQHASKQIADKQ | 50 |
| NT1p | YKGIIDCVVRIPKECGFLSFWRGNLANVIRYFPTQALNFAFKDKYKQIFL | 100 |
| NT2p | YKGIIDCVVRIPKECGVLSFWRCNLANVIRYFPTQALNFAFKDKYKQIFL | 100 |
| NT3p | YKGIIDCVVRIPKECGVLSFWRCNLANVIRYFPTQALNFAFKDKYKQIFL | 100 |
| NT1p | CGVDFHKKQFWRVYFAGNLASGGAAGATSLCFVYPLDFARTRLAADVGR-A | 149 |
| NT2p | CGVDKHTQFWLYFAGNLASGGAAGATSLCFVYPLDFARTRLAADVCKACA | 150 |
| NT3p | CGVDKHTQFWRYFAGNLASGGAAGATSLCFVYPLDFARTRLAADVCKSGT | 150 |
| NT1p | QREFHGLGDCITKTKKSDGIRGLYQGFNWSVQGIITYRAAYFGVYDTAKG | 199 |
| NT2p | EREFHGLGDCITKTKKSDGIRGLYQGFNWSVQGIITYRAAYFGVYDTAKG | 200 |
| NT3p | EREFHGLGDCIMKTKKSDGIRGLYQGFNWSVQGIITYRAAYFGVYDTAKG | 200 |
| NT1p | MLPDPKNIHIVSWMTAQSVTAVAGLISYPFDIVRRRMMMQSCKGADIM | 249 |
| NT2p | MLPDPKNIHIVSWMTAQSVTAVAGLISYPFDIVRRRMMMQSCKGADIM | 250 |
| NT3p | MLPDPKNIHIVSWMTAQSVTAVAGVLSYPFDIVRRRMMMQSCKGADIM | 250 |
| NT1p | YTGIVDCNRKIAFDEGCKAFFKQWNSWLRMGCAFVLVLVYDEIKKVT | 298 |
| NT2p | YTGIVDCNRKIAFDEGCKAFFKQWNSWLRMGCAFVLVLVYDEIKKVT | 299 |
| NT3p | YTGIVDCNRKIAFDEGCKAFFKQWNSWLRMGCAFVLVLVYDEIKKVT | 299 |

Figure 2

05810644-031401

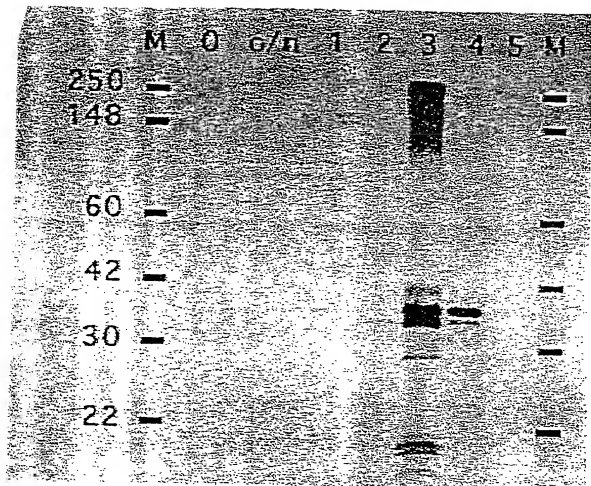


Figure 3

0810644.031401

M (+) 1 2 3 4

48 —

60 —

42 —

30 —

22 —

17 —



Figure 4

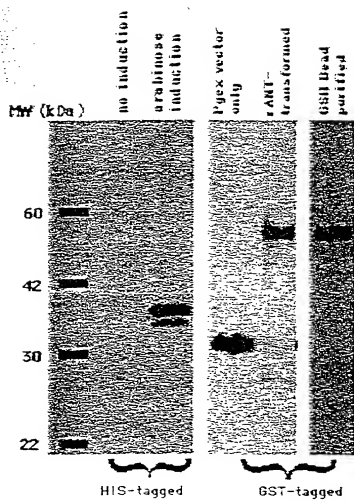


Figure 5

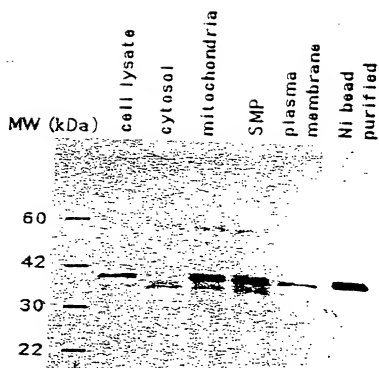


Figure 6

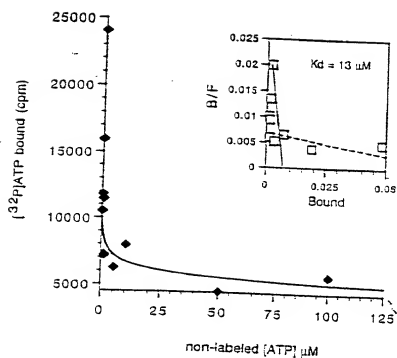


Figure 7

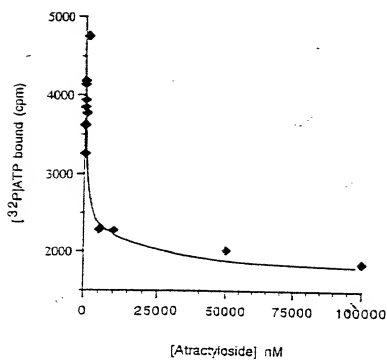


Figure 8

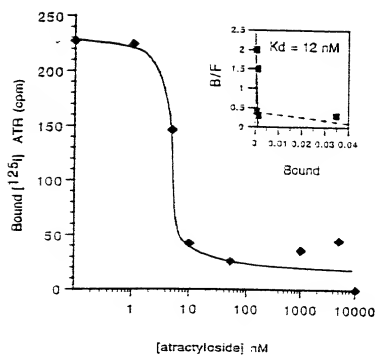


Figure 9

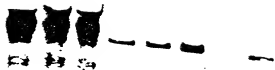


Figure 10 -

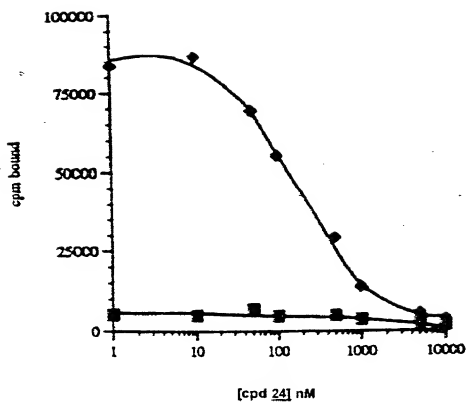


Figure 11

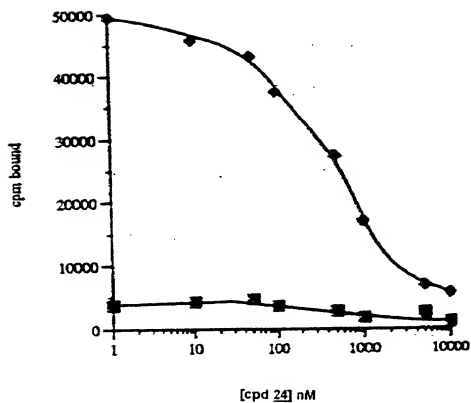


Figure 12

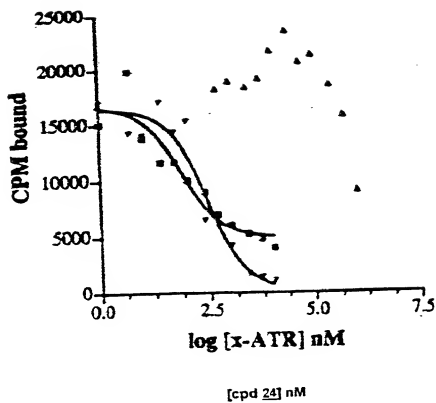


Figure 13

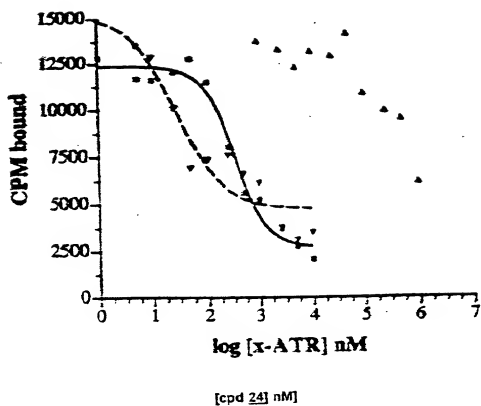


Figure 14

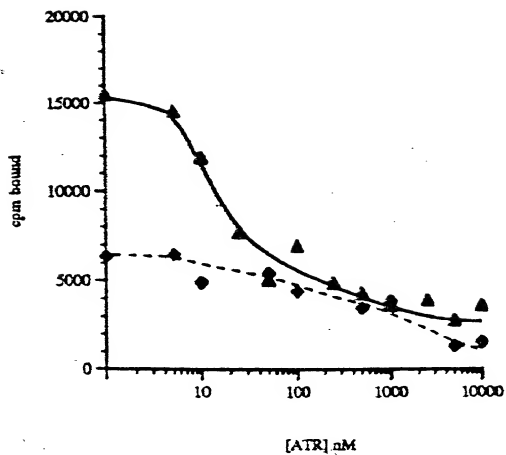


Figure 15

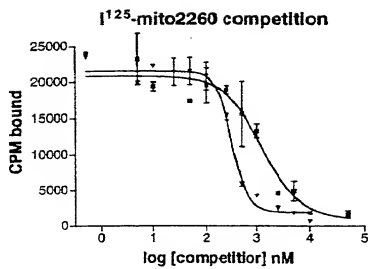


Figure 16

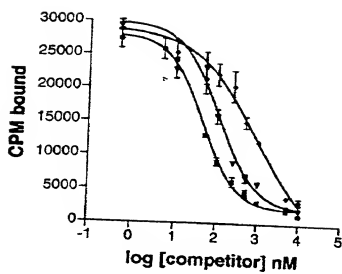


Figure 17

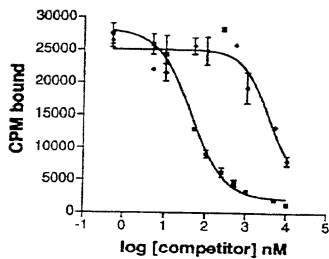


Figure 18

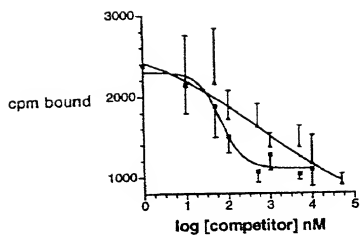


Figure 19